

# SEQUENCE LISTING

<110> OLSON, ERIC  
ARAI, AKIKO

<120> STARS - A MUSCLE-SPECIFIC ACTIN-BINDING PROTEIN

<130> MYOG:037US

<140> UNKNOWN

<141> 2003-08-20

<150> 60/404,706

<151> 2002-08-20

<160> 15

<170> PatentIn Ver. 2.1

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<222> (1)..(1146)

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Thr Gly Trp Leu Pro Gly Gly Thr Gln Asp Ser Pro Gln Ala Pro Lys	
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Pro Ile Thr Pro Pro Thr Ser His Gln Lys Ala Gln Ser Ala Pro Lys	
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Ser Pro Pro Arg Leu Pro Glu Gly His Gly Asp Gly Gln Ser Ser Glu	
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aaa gcc cct gag gtt tct cac atc aaa aag aaa gag gtg tcc aaa acg	336
Lys Ala Pro Glu Val Ser His Ile Lys Lys Lys Glu Val Ser Lys Thr	
100 105 110	

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Asp Ser Gly Tyr Gly Gly Glu Ala Glu Glu Arg Pro Glu Gln Asp Gly	
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Ile Gln Ser Gln Lys Leu Asn Pro Phe Ser Glu Glu Phe Asp Tyr Glu	
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Leu Ala Met Ser Thr Arg Leu His Lys Gly Asp Glu Gly Tyr Gly Arg	
275 280 285	
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290 295 300	
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His Ile Tyr Arg Glu Met Met Asp Met Cys Phe Ile Ile Cys Thr Met	
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 210 215 220  
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Thr Trp Lys Ser Asp Ser Val Asp Thr Glu Asp Ser Gly Tyr Gly Gly	
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Asp Met Glu Glu Arg Pro Glu Gln Asp Ala Ala Pro Val Ala Pro Ala	
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agg atc aaa cgc ccc ttg cac tcc cag gca aac agg tac tct gag cca	672
Arg Ile Lys Arg Pro Leu His Ser Gln Ala Asn Arg Tyr Ser Glu Pro	
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ctc aac tgt aag gcc cat cgg aaa tac agc caa gtg gac aac ttg aaa	720
Leu Asn Cys Lys Ala His Arg Lys Tyr Ser Gln Val Asp Asn Leu Lys	
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Asn Pro Phe Ser Asp Glu Phe Asp Tyr Asp Leu Ala Met Ser Thr Arg	
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ctc cac aag gga gac gag ggc tat ggc cgc ccc aaa gag gga agc aag	864
Leu His Lys Gly Asp Glu Gly Tyr Gly Arg Pro Lys Glu Gly Ser Lys	
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aca gct gaa agg gcc aag cga gcg gaa gag cac atc tat cgg gaa att	912
Thr Ala Glu Arg Ala Lys Arg Ala Glu Glu His Ile Tyr Arg Glu Ile	
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atg gaa ctg tgc ttt gtt atc cgc aca atg gct cgc cac aga cga gat	960
Met Glu Leu Cys Phe Val Ile Arg Thr Met Ala Arg His Arg Asp	
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Gly Lys Ile Gln Val Thr Phe Gly Glu Leu Phe Asp Arg Tyr Val Arg	
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Ile Ser Asp Lys Val Val Gly Ile Leu Met Arg Ala Arg Lys His Gly	
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1128

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 His Ile Lys Arg Lys Glu Val Thr Arg Thr Val Val Ser Lys Ala Tyr  
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 Gly Gly Val Ser Glu Ala Ile Gln Pro Glu Asn Asp Ile Asp Arg Ile  
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 Leu Leu Ser His Asp Ser Pro Thr Arg Arg Arg Lys Cys Thr Asn Leu  
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 Asp Met Glu Glu Arg Pro Glu Gln Asp Ala Ala Pro Val Ala Pro Ala  
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 Arg Ile Lys Arg Pro Leu His Ser Gln Ala Asn Arg Tyr Ser Glu Pro  
 210 215 220  
 Leu Asn Cys Lys Ala His Arg Lys Tyr Ser Gln Val Asp Asn Leu Lys  
 225 230 235 240  
 Gly Arg Trp Gln Gln Trp Ala Asp Glu His Val Gln Ser Gln Lys Leu  
 245 250 255  
 Asn Pro Phe Ser Asp Glu Phe Asp Tyr Asp Leu Ala Met Ser Thr Arg  
 260 265 270  
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 275 280 285  
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 290 295 300  
 Met Glu Leu Cys Phe Val Ile Arg Thr Met Ala Arg His Arg Arg Asp  
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 Gly Lys Ile Gln Val Thr Phe Gly Glu Leu Phe Asp Arg Tyr Val Arg  
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35 40 45  
Thr Arg Arg Arg Tyr Cys Gly Gly Lys Ala Gly Thr Phe Val Lys Ala  
50 55 60  
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65 70 75 80  
Asp Ala Asp Asp Ser Gly Leu Gly Glu Glu Ala Ser Leu Ser Asp Asn  
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Ser Asp Leu Asn Glu Asn Glu Pro Lys Lys His Val Asn Arg His Lys  
100 105 110  
Ile Lys Val Thr Thr Met Gly Asp Leu Arg Ser Arg Trp Gln Arg Phe  
115 120 125  
Ala Glu Asp His Met Glu Gly Gln Lys Leu Asn Pro Phe Ser Glu Glu  
130 135 140

Phe Asp Tyr Asp His Ala Met Ala Thr Arg Leu His Lys Gly Asp Ala  
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 Glu Met Glu Gln Asn Val Ala Thr Gln Ser Lys Asp Asp Val Tyr Ser  
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 aaa gat ttt act caa aag aaa atg gac aag tcc agt agc gaa tat gga 144  
 Lys Asp Phe Thr Gln Lys Lys Met Asp Lys Ser Ser Ser Glu Tyr Gly  
 35 40 45  
 cgg cca aaa cca gga act ctt aca gag caa aga gct aaa aaa gct gcc 192  
 Arg Pro Lys Pro Gly Thr Leu Thr Glu Gln Arg Ala Lys Lys Ala Ala  
 50 55 60  
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 65 70 75 80  
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 Tyr Gly Lys Gln Glu Lys Glu Gly Asp Pro Ile Arg Ile Thr Phe Gly  
 85 90 95  
 aga ctt ttc aca att tat gtc aat att tct gat aag gta gtt gga acc 336  
 Arg Leu Phe Thr Ile Tyr Val Asn Ile Ser Asp Lys Val Val Gly Thr  
 100 105 110  
 ctt ttg cga gct cgt aaa cac aaa atg ata gat ttt gaa gga gaa atg 384  
 Leu Leu Arg Ala Arg Lys His Lys Met Ile Asp Phe Glu Gly Glu Met  
 115 120 125  
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 Leu Phe Gln Lys Arg Asp Asp His Val Ile Ile Thr Leu Leu Leu Ser



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Arg Leu Phe Thr Ile Tyr Val Asn Ile Ser Asp Lys Val Val Gly Thr	
100 105 110	
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115 120 125	
Leu Phe Gln Lys Arg Asp Asp His Val Ile Ile Thr Leu Leu Leu Ser	
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 <213> *Drosophila melanogaster*

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Gln Asp Gly Arg Ala Ala Ser Pro Lys Pro Thr Phe Ser Lys Asp Gln	
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Tyr	Gly	Lys	Pro	Leu	Ala	Gly	Ser	Leu	Thr	Glu	Met	Arg	Gly	Gln	Lys	65	70	75	80
Ala	Asn	Ile	His	Val	Met	Lys	Glu	Met	Leu	Glu	Leu	Cys	Gln	Ile	Ile	85	90	95	
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His	Lys	Leu	Val	Asp	Phe	Glu	Gly	Glu	Met	Leu	Tyr	Gln	Arg	Arg	Asp	165	170	175	
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Glu	Met	Glu	Ala	Lys	Ile	Glu	Asp	Ile	Lys	Arg	Ala	Ala	Ser	Pro	Ala	195	200	205	
Pro	Pro	Gln	Ser	Thr	Ser	Val	Leu	Met	Asp	Arg	Ser	Ala	His	Glu	Gln	210	215	220	
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Ser	Lys	Ser	Ala	Ser	Pro	Ala	Pro	Lys	Ala	Pro	Val	Pro	Val	Pro	Ala	245	250	255	
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Val	Thr	Glu	Ala	Pro	Ser	Thr	Glu	Glu	Thr	Thr	Pro	Thr	Thr	Ser	Glu	325	330	335	
Pro	Gln	Ala	Glu	Glu	Ala	Pro	Ala	Ala	Val	Ala	Pro	Ala	Gly	Pro	Ala	340	345	350	
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Primer

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